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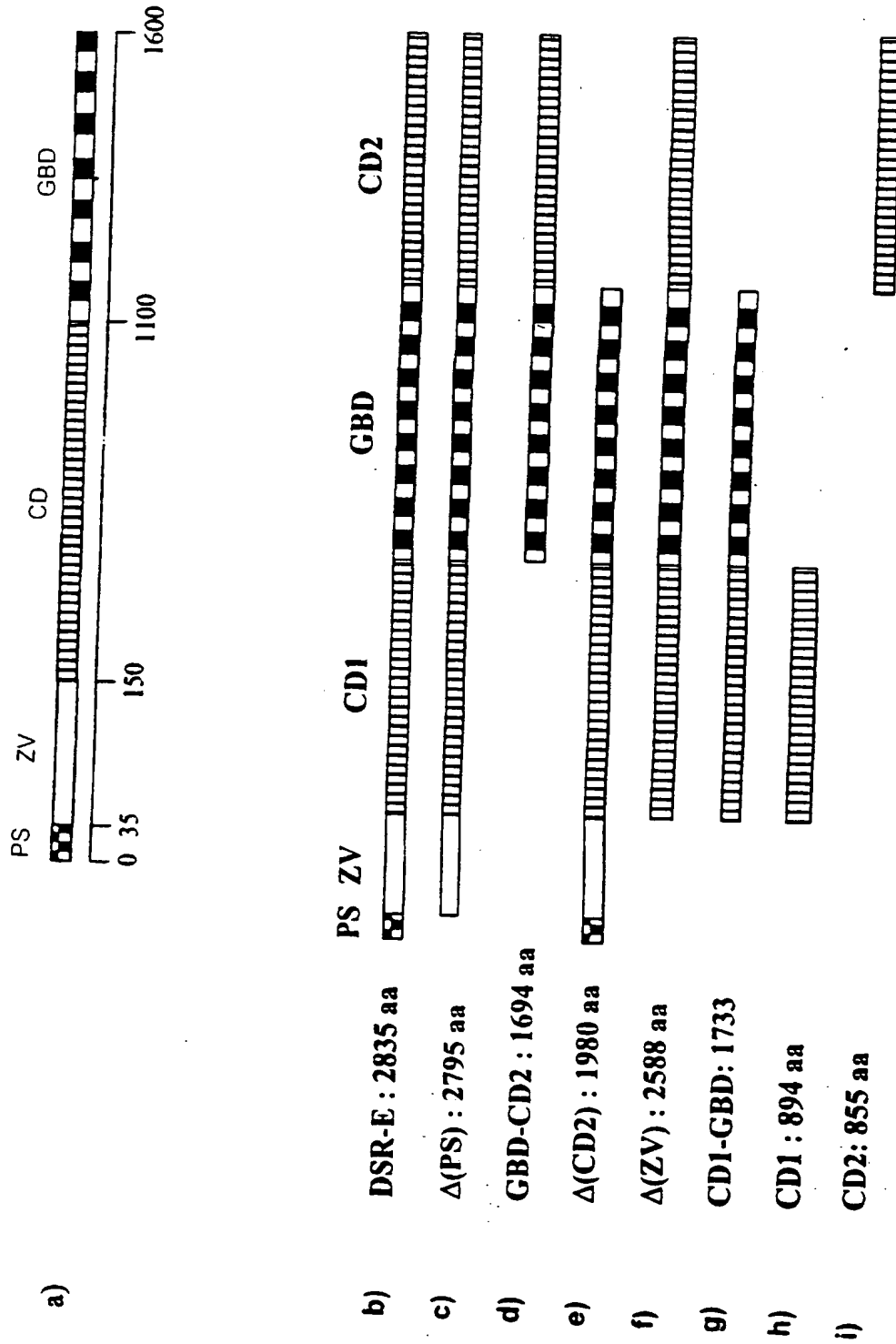


Fig. 1

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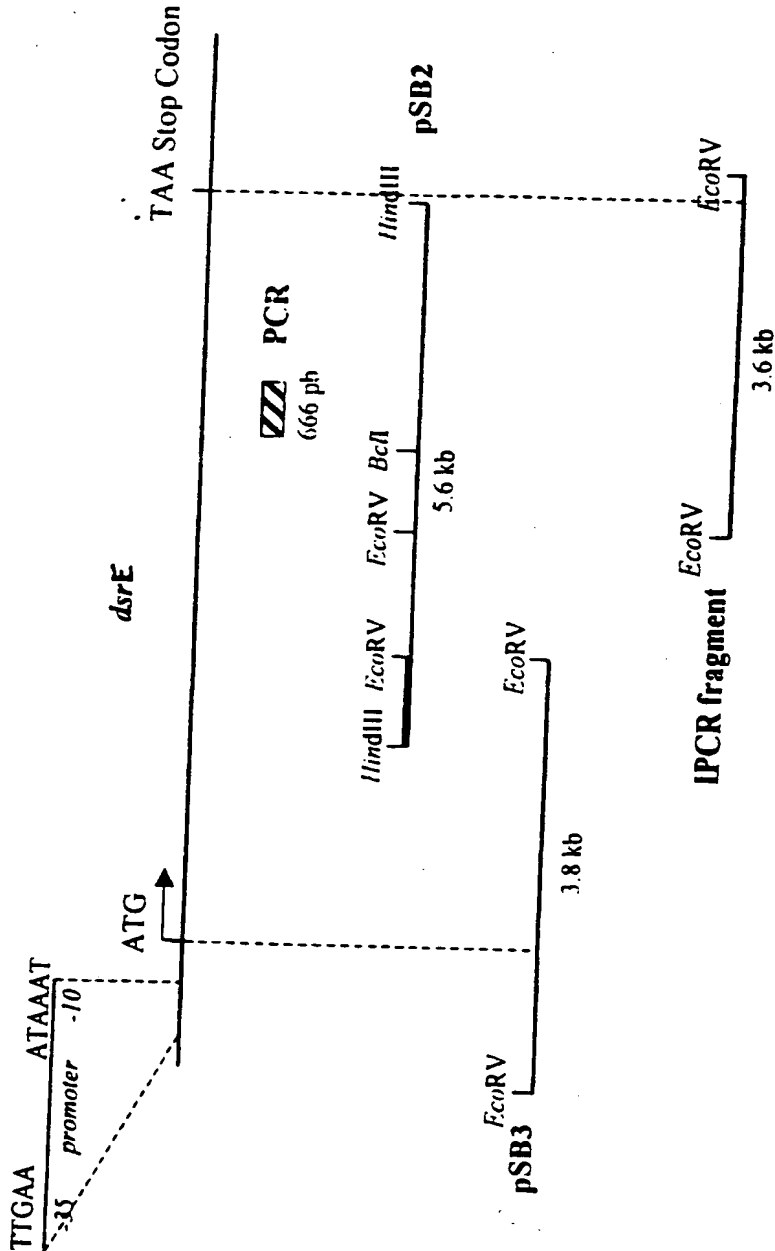


Fig. 2

DSR-E	MRDMRVICDRKKLYKSGKVLVTAG-IFALMMEGVTTASVSA
DSR-B	MEMIKERNVRRKLYKSGKSWVIGGLILSTIMLSMTATS---
DSR-S	-MPFTEKVMRRKLYKVGKSWVVG-----VCAFALTAS-----
ASR	-MKQQTETVTRK-YKSGKVVAAATAFAVLGVSTVTTVHA-

Fig. 3

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⁷³AAKVAVATTP-AT
⁸⁶PVADKT VSA
⁹⁵PAADKAVDTTSSTT
¹⁰⁹PATDKAVDTTP-TT
¹²²PAADKAVDTTP-TT
¹³⁵PAADKAVDTTP-TT
¹⁴⁸PAANKAVDTTP-AT
¹⁶¹AATDKAV-ATP-AT
¹⁷³PAADKLANTT--AT
¹⁸⁵----DKAVATTP-AT
¹⁹⁶PVANKAA

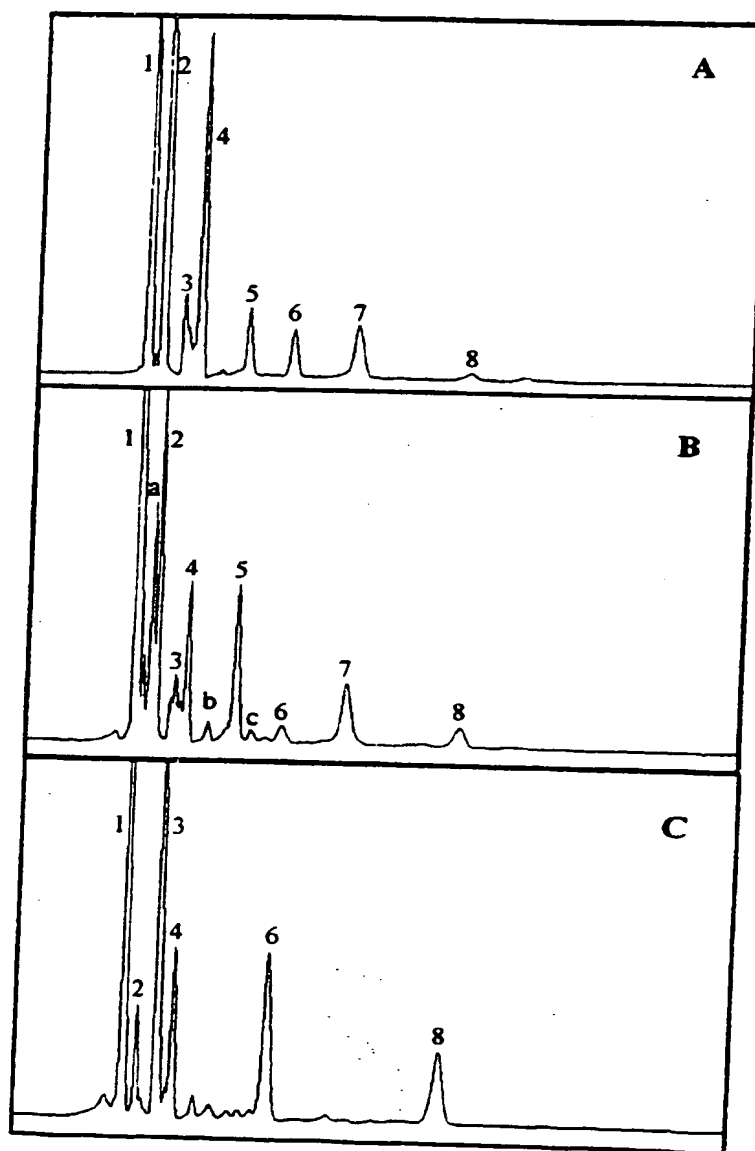
 PAADKAVDTTP-~~A~~/T ← Proposed consensus sequence for S repeat

Fig. 4

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Fig. 6



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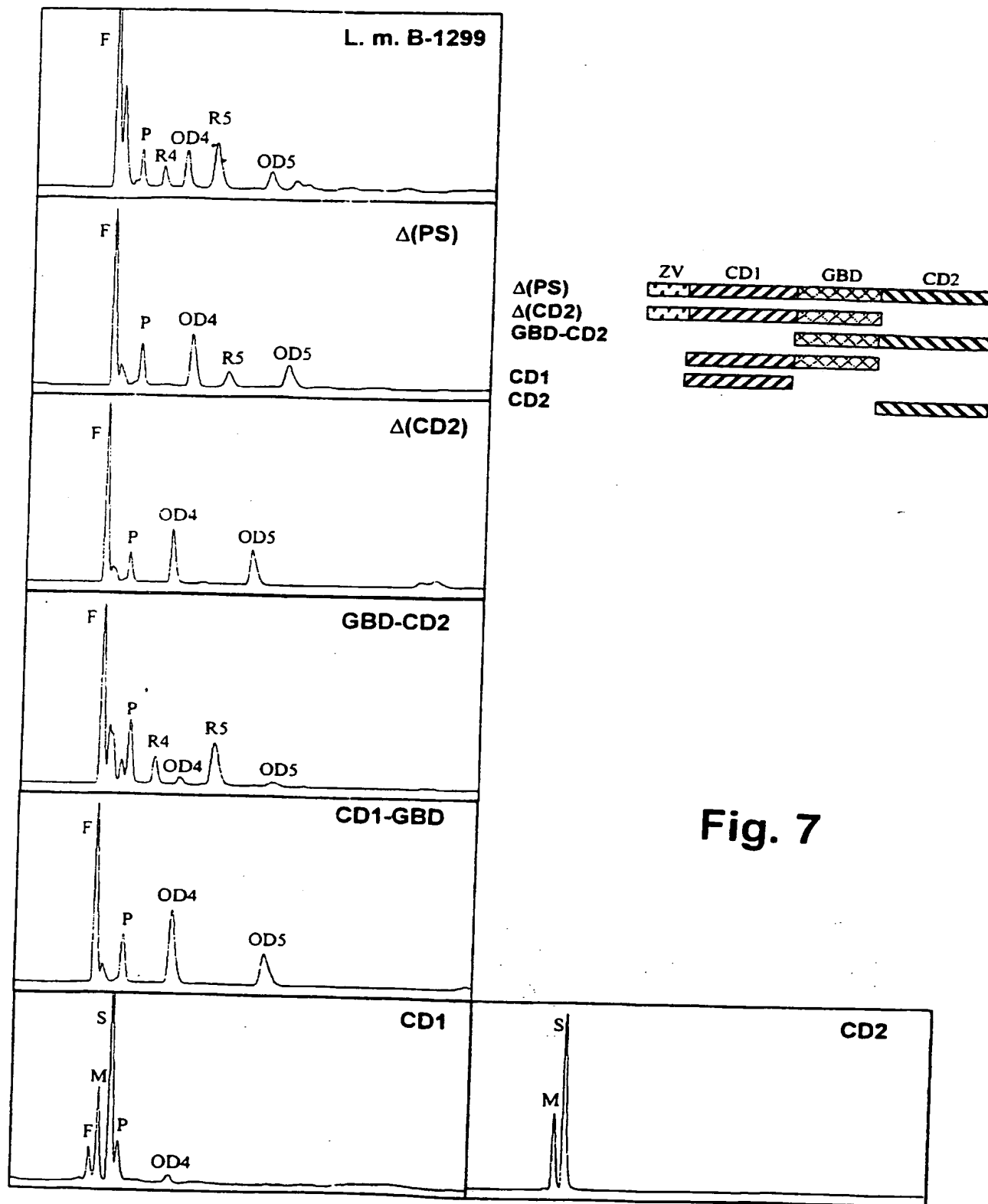


Fig. 7